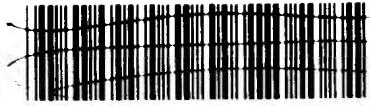


Raw Sequence Listing Error Summary

<u>ERROR DETECTED</u>	<u>SUGGESTED CORRECTION</u>	<u>SERIAL NUMBER:</u> <u>09/804,733B</u>
ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE		
1 <input type="checkbox"/> Wrapped Nucleic Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."	
2 <input type="checkbox"/> Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.	
3 <input type="checkbox"/> Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.	
4 <input type="checkbox"/> Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.	
5 <input type="checkbox"/> Variable Length	Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.	
6 <input type="checkbox"/> PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.	
7 <input type="checkbox"/> Skipped Sequences (OLD RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped. Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.	
8 <input type="checkbox"/> Skipped Sequences (NEW RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000	
9 <input type="checkbox"/> Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.	
10 <input type="checkbox"/> Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence	
11 <input type="checkbox"/> Use of <220>	Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)	
12 <input type="checkbox"/> PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.	
13 <input type="checkbox"/> Misuse of n	n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.	



1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/804,733B

DATE: 12/18/2002

TIME: 11:01:59

Input Set : A:\MONS016US.TXT

Output Set: N:\CRF4\12182002\I804733B.raw

3 <110> APPLICANT: WANG et al.
 5 <120> TITLE OF INVENTION: RECOMBINANT PROTEINS CONTAINING REPEATING UNITS
 7 <130> FILE PREFERENCE: MONS:016US
 9 <140> CURRENT APPLICATION NUMBER: 09/804,733B
 10 <141> CURRENT FILING DATE: 2001-03-13
 11 <150> PRIORITY APPLICATION NUMBER: 60/188,990
 13 <151> PRIORITY FILING DATE: 2000-03-13
 15 <160> NUMBER OF SEQ ID NOS: 29
 17 <170> SOFTWARE: PatentIn version 3.0
 18 <110> SEQ ID NO: 1
 19 <211> LENGTH: 5
 20 <212> TYPE: PPT
 21 <213> ORGANISM: Euthynnus pelamis
 22 <220> FEATURE:
 23 <211> NAME/KEY: PEPTIDE
 24 <211> LOCATION: (1)..(5)
 25 <400> SEQUENCE: 1
 26 Lys Pro Asn Met
 27 1 5
 28 <210> SEQ ID NO: 2
 29 <211> LENGTH: 4
 30 <212> TYPE: PPT
 31 <213> ORGANISM: Euthynnus pelamis
 32 <220> FEATURE:
 33 <211> NAME/KEY: PEPTIDE
 34 <211> LOCATION: (1)..(4)
 35 <400> SEQUENCE: 2
 36 Lys Pro Asn Met
 37 1
 38 <210> SEQ ID NO: 3
 39 <211> LENGTH: 4
 40 <212> TYPE: PPT
 41 <213> ORGANISM: Euthynnus pelamis
 42 <220> FEATURE:
 43 <211> NAME/KEY: PEPTIDE
 44 <211> LOCATION: (1)..(4)
 45 <400> SEQUENCE: 3
 46 Val Val Tyr Pro
 47 1
 48 <210> SEQ ID NO: 4
 49 <211> LENGTH: 15
 50 <212> TYPE: DNA
 C--> 64 <213> ORGANISM: Artificial/Unknown

Does Not Comply
 Corrected Diskette Needed

ppr 1-6

invalid response - see item 10 on
 Error Summary
 Sheet

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/804,733B

DATE: 11/16/02
TIME: 10:11:14

Input Set: A:\MONS016US.TXT
Output Set: N:\CRF4\12182002\I804733B.raw

66 <210> FEATURE:
67 <210> NAME/KEY: misc_feature
68 <210> LOCATION: (1)..(15)
69 <210> OTHER INFORMATION: Degenerate sequence
72 <210> FEATURE:
73 <210> NAME/KEY: misc_feature
74 <210> LOCATION: (1)..(15)
75 <210> OTHER INFORMATION: n=a, t, c or g; r=a or g; y=c or t
78 <400> SEQUENCE: 4

W16 79 ctnaarccna ayatg

15

82 <400> SEQ ID NO: 5
83 <400> LENGTH: 60
84 <400> TYPE: DNA

C--> 85 <213> ORGANISM: Artificial/Unknown

87 <210> FEATURE:
88 <210> NAME/KEY: misc_feature
89 <210> LOCATION: 1)..(60)
90 <210> OTHER INFORMATION: n=any nucleotide; r=a or g; y=c or t
93 <210> FEATURE:
94 <210> NAME/KEY: misc_feature
95 <210> LOCATION: (1)..(60)
96 <210> OTHER INFORMATION: Degenerate sequence
99 <400> SEQUENCE: 5

W16 -> 100 ctnaarccna ayatgctnaa rccnaayatg ctnaarccna ayatgctnaa rccnaayatg

60

103 <400> SEQ ID NO: 6
104 <400> LENGTH: 60
105 <400> TYPE: DNA

C--> 106 <213> ORGANISM: Artificial/Unknown

108 <210> FEATURE:
109 <210> NAME/KEY: misc_feature
110 <210> LOCATION: (1)..(60)
111 <210> OTHER INFORMATION: n=any nucleotide, r=a or g, y=c or t
114 <210> FEATURE:
115 <210> NAME/KEY: misc_feature
116 <210> LOCATION: (1)..(60)
117 <210> OTHER INFORMATION: degenerate sequence
120 <400> SEQUENCE: 6

W16 -> 121 catrttnggy ttnagcatrt tnggyttnag catrttnggy ttnagcatrt tnggyttnag

60

124 <400> SEQ ID NO: 7
125 <400> LENGTH: 25
126 <400> TYPE: DNA

C--> 127 <213> ORGANISM: Artificial/Unknown

129 <210> FEATURE:
130 <210> NAME/KEY: misc_feature
131 <210> LOCATION: (1)..(25)
132 <210> OTHER INFORMATION: Primer
134 <210> FEATURE:
135 <210> NAME/KEY: misc_feature
137 <210> LOCATION: (1)..(25)

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/804,733B

DATE: 10/17/02
TIME: 11:15

Input Set : A:\MONS016US.TXT
Output Set: N:\CRF4\12182002\I804733B.raw

137 <213> OTHER INFORMATION: n=any nucleotide; r=a or g; y=c or t
141 <400> SEQUENCE: 7

OK-> 142 aaagaattcc tnaarccnaa yatgc 25
143 <210> SEQ ID NO: 8
144 <211> LENGTH: 27
145 <212> TYPE: DNA

C--> 148 <213> ORGANISM: Artificial/Unknown

150 <220> FEATURE:
151 <221> NAME/KEY: misc_feature
152 <222> LOCATION: (1)..(27)
153 <223> OTHER INFORMATION: Primer
154 <224> FEATURE:
155 <225> NAME/KEY: misc_feature
156 <226> LOCATION: (1)..(27)
157 <213> OTHER INFORMATION: n=any nucleotide; r=a or g; y=c or t

161 <400> SEQUENCE: 8

OK-> 163 aaagcggccg ccatrtnnng yttnagc 27

166 <210> SEQ ID NO: 9
167 <211> LENGTH: 36
168 <212> TYPE: DNA

C--> 169 <213> ORGANISM: Artificial/Unknown

171 <220> FEATURE:
172 <221> NAME/KEY: misc_feature
173 <222> LOCATION: (1)..(20)
174 <223> OTHER INFORMATION: Primer
175 <224> SEQUENCE: 9
176 tcatttttgtt cactatagg 20
180 <210> SEQ ID NO: 10
181 <211> LENGTH: 19
182 <212> TYPE: DNA

C--> 184 <213> ORGANISM: Artificial/Unknown

187 <220> FEATURE:
188 <221> NAME/KEY: misc_feature
189 <222> LOCATION: (1)..(19)
190 <223> OTHER INFORMATION: Primer
191 <224> SEQUENCE: 10
192 tcatttttgtt cggatcgcc 19
196 <210> SEQ ID NO: 11
197 <211> LENGTH: 47
198 <212> TYPE: DNA

C--> 199 <213> ORGANISM: Artificial/Unknown

201 <220> FEATURE:
202 <221> NAME/KEY: misc_feature
203 <222> LOCATION: (1)..(48)
204 <213> OTHER INFORMATION: n=any nucleotide; y=c or t
205 <224> FEATURE:
206 <225> NAME/KEY: misc_feature
207 <226> LOCATION: (1)..(48)
210 <223> OTHER INFORMATION: Degenerate sequence

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/804,733B

DATE: 12/18/02
TIME: 10: 07: 47

Input Set : A:\MONS016US.TXT
Output Set: N:\CRF4\12182002\I804733B.raw

W 213 <400> SEQUENCE: 11 48
W-> 214 gtngtntayc cngtngtnta yccngtngtn tayccngtng tntayccn
215 <100> SEQ ID NO: 12
216 <100> LENGTH: 48
217 <100> TYPE: DNA
C--> 220 <213> ORGANISM: Artificial/Unknown
221 <100> FEATURE:
222 <100> NAME/KEY: misc_feature
223 <100> LOCATION: (1)..(48)
224 <100> OTHER INFORMATION: n=any nucleotide; r=a or g
225 <100> FEATURE:
226 <100> NAME/KEY: misc_feature
227 <100> LOCATION: (1)..(48)
228 <100> OTHER INFORMATION: Degenerate sequence
229 <400> SEQUENCE: 12
W 235 nggrtanaacn acnggrtana cnacnggrta nacnaacnggr tanacnac 48
236 <100> SEQ ID NO: 13
237 <100> LENGTH: 31
238 <100> TYPE: DNA
C--> 241 <213> ORGANISM: Artificial/Unknown
242 <100> FEATURE:
243 <100> NAME/KEY: misc_feature
244 <100> LOCATION: (1)..(31)
245 <100> OTHER INFORMATION: Forward primer
246 <100> FEATURE:
247 <100> NAME/KEY: misc_feature
248 <100> LOCATION: (1)..(31)
249 <100> OTHER INFORMATION: n=any nucleotide; y=c or t
250 <400> SEQUENCE: 13
W-> 256 aaaggatccg tngtntaycc ngtngtntay ccn 33
257 <100> SEQ ID NO: 14
258 <100> LENGTH: 31
259 <100> TYPE: DNA
C--> 262 <213> ORGANISM: Artificial/Unknown
263 <100> FEATURE:
264 <100> NAME/KEY: misc_feature
265 <100> LOCATION: (1)..(31)
266 <100> OTHER INFORMATION: Reverse primer
267 <100> FEATURE:
268 <100> NAME/KEY: misc_feature
269 <100> LOCATION: (1)..(31)
270 <100> OTHER INFORMATION: n=any nucleotide; r=a or g
271 <400> SEQUENCE: 14
W-> 277 cccaagcttn ggrtanacna cnggrrtanac nac 33
278 <100> SEQ ID NO: 15
279 <100> LENGTH: 41
280 <100> TYPE: DNA
C--> 283 <213> ORGANISM: Artificial/Unknown
284 <100> FEATURE:

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/804,733B

DATE: 12/18/02
 TIME: 12: 1:54

Input Set : A:\MONS016US.TXT
 Output Set: N:\CRF4\12182002\I804733B.raw

286 <1..> NAME/KEY: misc_feature
 287 <1..> LOCATION: (1)..(45)
 288 <1..> OTHER INFORMATION: n=any nucleotide
 289 <1..> FEATURE:
 290 <1..> NAME/KEY: misc_feature
 291 <1..> LOCATION: (1)..(45)
 292 <1..> OTHER INFORMATION: Degenerate sequence
 297 <4..> SEQUENCE: 16

W-> 298 gtnccnccng tnccnccngt nccnccngtn ccnccngtnc cncn 45

301 <1..> SEQ ID NO: 16

302 <1..> LENGTH: 45

303 <1..> TYPE: DNA

C--> 304 <213> ORGANISM: Artificial/Unknown

310 <1..> FEATURE:

311 <1..> NAME/KEY: misc_feature

312 <1..> LOCATION: (1)..(45)

313 <1..> OTHER INFORMATION: n=any nucleotide

314 <1..> FEATURE:

315 <1..> NAME/KEY: misc_feature

316 <1..> LOCATION: (1)..(45)

317 <1..> OTHER INFORMATION: Degenerate sequence

318 <1..> SEQUENCE: 16

W-> 319 ngngggnacn ggnggnacng gnggnacnngg ngnacnnggn ggnac 45

320 <1..> SEQ ID NO: 17

321 <1..> LENGTH: 36

322 <1..> TYPE: DNA

C--> 325 <213> ORGANISM: Artificial/Unknown

323 <1..> FEATURE:

324 <1..> NAME/KEY: misc_feature

325 <1..> LOCATION: (1)..(36)

326 <1..> OTHER INFORMATION: Forward primer

327 <1..> FEATURE:

328 <1..> NAME/KEY: misc_feature

329 <1..> LOCATION: (1)..(36)

330 <1..> OTHER INFORMATION: n=any nucleotide

331 <1..> SEQUENCE: 17

W-> 340 aaaggatccg tnccnccngt nccnccngtn ccnccn 36

341 <1..> SEQ ID NO: 18

342 <1..> LENGTH: 36

343 <1..> TYPE: DNA

C--> 346 <213> ORGANISM: Artificial/Unknown

344 <1..> FEATURE:

345 <1..> NAME/KEY: misc_feature

346 <1..> LOCATION: (1)..(36)

347 <1..> OTHER INFORMATION: Reverse primer

348 <1..> FEATURE:

349 <1..> NAME/KEY: misc_feature

350 <1..> LOCATION: (1)..(36)

351 <1..> OTHER INFORMATION: n=any nucleotide

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/804,733B

DATE: 12/18/02
TIME: 11:45:11

Input Set : A:\MONS016US.TXT
Output Set: N:\CRF4\12182002\I804733B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:4; N Pos. 3,9
Seq#:5; N Pos. 3,9,18,24,33,39,46,54
Seq#:6; N Pos. 7,13,22,28,37,43,52,58
Seq#:7; N Pos. 12,18
Seq#:8; N Pos. 18,24
Seq#:11; N Pos. 3,6,12,15,18,24,27,30,36,39,42,48
Seq#:12; N Pos. 1,7,10,13,19,22,25,31,34,37,43,46
Seq#:13; N Pos. 12,15,21,24,27,33
Seq#:14; N Pos. 10,16,19,22,23,31
Seq#:15; N Pos. 3,6,9,12,15,16,21,24,27,30,33,36,39,42,45
Seq#:16; N Pos. 1,4,7,10,13,16,18,22,25,28,31,34,37,40,43
Seq#:17; N Pos. 12,15,18,21,24,27,30,33,36
Seq#:18; N Pos. 10,15,16,19,22,25,28,31,34
Seq#:20; N Pos. 3,6,9,12,15,27,31,33,36,42
Seq#:21; N Pos. 7,13,16,19,22,31,37,40,43,46
Seq#:22; N Pos. 13,16,19,22,28,31,40,43,46,52
Seq#:23; N Pos. 19,25,28,31,34,40,43,46,55,61,64,67,70,76,79,82
Seq#:24; N Pos. 3,6,9,15,18,21,24,30,39,42,45,51,54,57,60,66
Seq#:25; N Pos. 7,13,16,19,22,25,31,34,43,49,52,55,58,64,67,70
Seq#:26; N Pos. 13,16,19,25,28,31,34,40,49,52,55,61,64,67,70,76
Seq#:27; N Pos. 19,25,28,31,34,40,43,46,55,61,64,67,70,76,79,82
Seq#:28; N Pos. 19,25,28,31,34,40,43,46,55,61,64,67,70,76,79,82

VERIFICATION SUMMARY

PATIENT APPLICATIONN: US/09/804,733B

PAGE: 11 OF 11

TIME: 11:11:

Input Set : A:\MONS016US.TXT

Output Set: N:\CRF4\12182002\I804733B.raw

L:64 M:220 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:4
L:79 M:341 W: (4) "n" or "Kaa" used, for SEQ ID#:4 after pos.:4
L:85 M:211 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:5
L:100 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:5 after pos.:6
L:106 M:11 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:6
L:111 M:341 W: (4) "n" or "Kaa" used, for SEQ ID#:6 after pos.:6
L:117 M:11 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:7
L:142 M:341 W: (4) "n" or "Kaa" used, for SEQ ID#:7 after pos.:7
L:148 M:11 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:8
L:163 M:341 W: (4) "n" or "Kaa" used, for SEQ ID#:8 after pos.:8
L:179 M:11 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:9
L:184 M:11 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:10
L:199 M:341 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:11
L:211 M:341 W: (4) "n" or "Kaa" used, for SEQ ID#:11 after pos.:8
L:213 M:11 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:12
L:215 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:12 after pos.:8
L:241 M:341 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:13
L:256 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:13 after pos.:8
L:262 M:341 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:14
L:278 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:14 after pos.:8
L:284 M:341 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:15
L:299 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:15 after pos.:8
L:314 M:341 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:16
L:329 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:16 after pos.:8
L:335 M:341 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:17
L:346 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:17 after pos.:8
L:356 M:341 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:18
L:361 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:18 after pos.:8
L:367 M:341 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:19
L:382 M:341 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:20
L:398 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:20 after pos.:8
L:404 M:341 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:21
L:419 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:21 after pos.:8
L:434 M:341 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:22
L:445 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:22 after pos.:8
L:446 M:341 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:23
L:461 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:23 after pos.:8
L:466 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:24 after pos.:8
L:481 M:341 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:24
L:486 M:341 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:25
L:501 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:25 after pos.:8
L:503 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:25 after pos.:8
L:505 M:341 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:26
L:513 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:26 after pos.:8
L:515 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:26 after pos.:8
L:531 M:341 C: Keyword misspelled or invalid format, -213- ORGANISM for SEQ ID#:27
L:546 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:27 after pos.:8
L:548 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:27 after pos.:8

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/804,733B

DATE: 11/17/2009

TIME: 11:11:11

Input Set: A:\MONS016US.TXT

Output Set: N:\CRF4\12182002\I804733B.raw

L:634 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:17
L:669 M:341 W: 146 "n" or "Xaa" used, for SEQ ID#:28 after pos.:1
L:671 M:341 W: 146 "n" or "Xaa" used, for SEQ ID#:28 after pos.:66
L:677 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:19